



Figure S2. Hairpin secondary structures formed by the direct repeats from each CRISPR found in coagulase negative *Staphylococcus* studied in this work. Structures were predicted by RNAfold. Reference: Gruber AR, Lorenz R, Bernhart SH, Neuböck R, Hofacker IL. (2008). The Vienna RNA Websuite. Nucleic Acids Res. 36 (2): W70-W74.